

Advanced Regression: Linear and generalised linear models II

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Generalised linear model

- Basic definition
- Technical details on exponential families and GLMs
- Logistic regression and binary outcomes
- Generalised linear models in R

Generalised linear model (GLM)

- Linear models can only model a quantitative outcome.
- Quantitative outcomes are defined as a real number, taking possible values from $-\infty$ to $+\infty$.
- Many important data types can by definition not be modelled using a linear model:
 - Dichotomous or binary \rightarrow only takes two values, 0 or 1
 - Counts \rightarrow only positive integers (0,1,2,3,...)

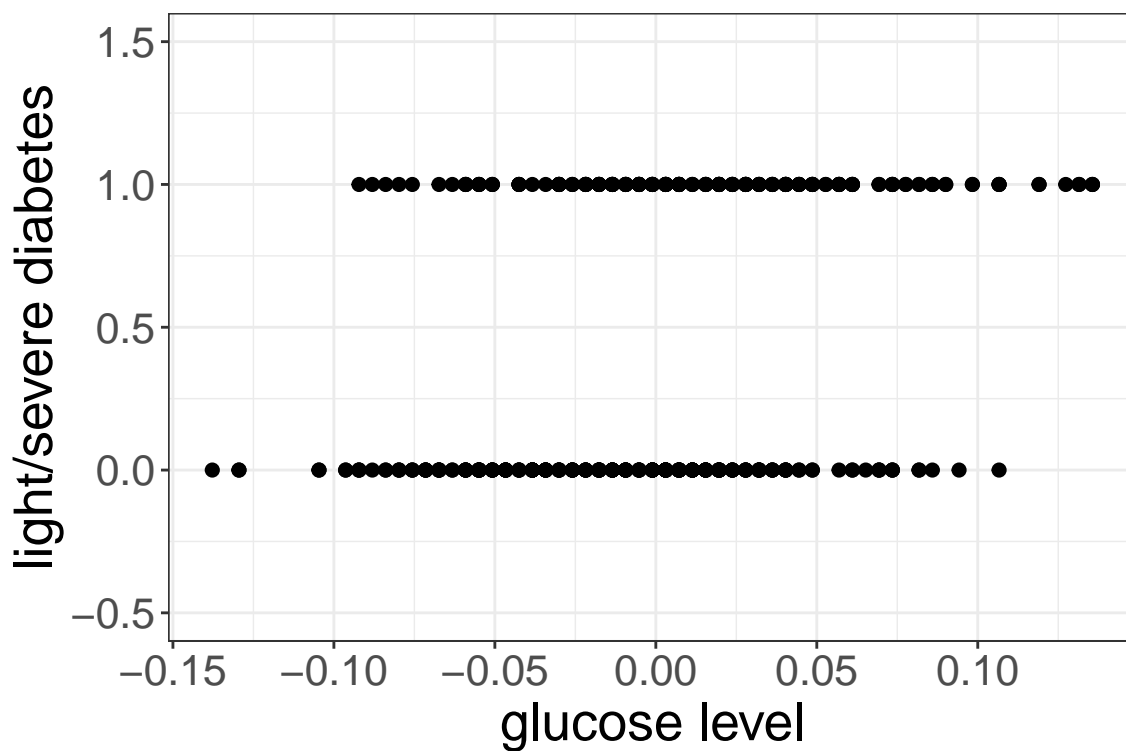
Note

Flexible generalization of ordinary linear regression that allows for response variables that have error distribution models other than a normal distribution. Residuals are an important quantity for model diagnostics.

Binary outcome and logistic regression

Example: Case-control study

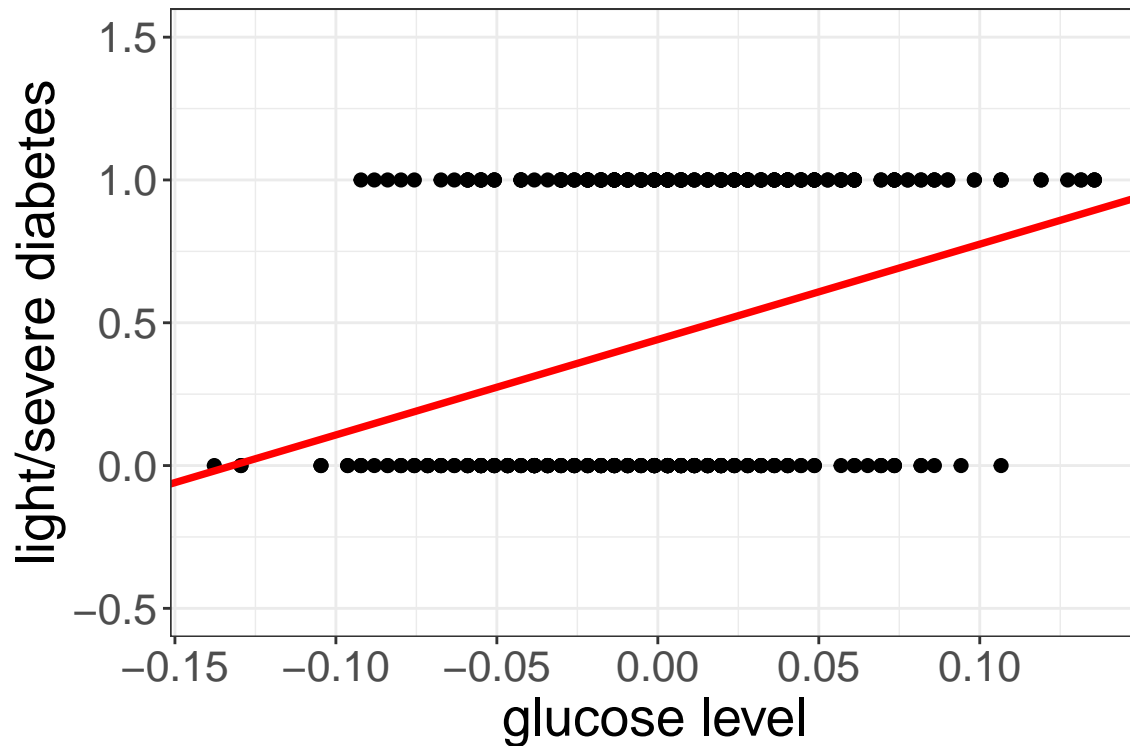
$$y_i = \begin{cases} 1, & \text{If subject } i \text{ is a case} \\ 0, & \text{If subject } i \text{ is a control} \end{cases}$$



Binary outcome and logistic regression

$$y = \underbrace{\alpha + \beta x}_{\text{Linear predictor}} + \epsilon$$

- Linear predictor: $\eta = \alpha + \beta x$ is defined from $-\infty$ to $+\infty$.
- But y only 0 or 1 \rightarrow The linear regression do not match the data well.



How should we model this data?

Key idea 1: Instead of modelling the outcomes ($y = 0$ or $y = 1$) directly, logistic regression models the probability for $y = 1$ denotes as

- $P(y = 1 \mid x)$

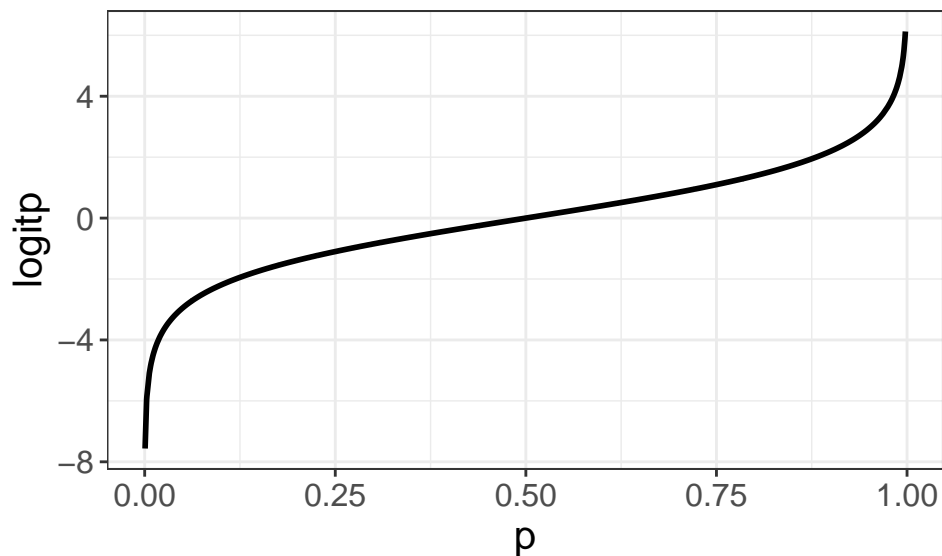
Notes on probabilities for binary data:

- Probabilities can take values from 0 to 1
- Probabilities are symmetric: $P(y = 1 \mid x) = 1 - P(y = 0 \mid x)$

How should we model this data?

Key idea 2: Transform the linear predictor $\eta = \alpha + \beta x_i$ (quantitative, can take values from $-\infty$ to ∞) to lie in the Interval $[0, 1]$, which is valid for probabilities.

This can be achieved using the logit function: $\text{logit}(p) = \log(p/(1 - p))$



Logistic regression

$$\text{logit}(P(y = 1 \mid x)) = \log(P(y = 1 \mid x) / (1 - P(y = 1 \mid x))) = \alpha + \beta x$$

- **Interpretation:** The regression coefficient β in logistic regression represents the **log odds ratio** between $y = 0$ and $y = 1$.
- **Estimation:** Maximum likelihood

Technical details

- Many important outcome types can be accommodated by GLMs.
- Each of these distributions has a location parameter, e.g. μ for the Gaussian, p for the Bernoulli and Binomial.
- The natural link function between the location parameter and the linear predictor can be derived from the mathematical form of the distribution.

Response	Distribution	E(y)	Link (g)
Continuous	Gaussian	μ	1 (identity)
Dichotomous	Bernoulli	p	logit
Counts	Poisson	λ	log

https://en.wikipedia.org/wiki/Generalized_linear_model

Technical details: GLM

The GLM consists of three elements:

1. A probability distribution from the exponential family. Note: Only distributions that can be formulated as an exponential family can be modelled as GLM.
2. A linear predictor $\eta = x\beta$
3. A link function g such that $E(y) = \mu = g^{-1}(\eta)$

Technical details: Exponential families

An exponential family is a set of probability distributions of the following form:

$$f_x(x | \theta) = h(x) \exp\{\eta(\theta) \times T(x) - A(\theta)\}$$

where

- θ is the parameter of interest.
- $T(x)$ is a sufficient statistic.
- $\eta(\theta)$ is the natural parameter or link function.

Gaussian distribution as exponential distribution

Gaussian distribution with unknown μ , but known σ :

$$f_\sigma(x | \mu) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left\{-\frac{(x - \mu)^2}{2\sigma^2}\right\} = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left\{-\frac{x^2}{2\sigma^2} + \frac{x\mu}{\sigma^2} - \frac{\mu^2}{2\sigma^2}\right\}$$

- $\theta = \mu$
- $h(x) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left\{-\frac{x^2}{2\sigma^2}\right\}$
- $T(x) = \frac{x}{\sigma}$
- $\eta(\mu) = \frac{\mu}{\sigma}$
- $A(\mu) = \frac{\mu^2}{2\sigma^2}$

Logistic regression and binary outcomes

Binomial distribution with known number of trials n , but unknown probability p :

$$f(x | p) = \binom{n}{x} p^x (1-p)^{n-x} = \quad (1)$$

$$= \binom{n}{x} \exp\{x \log(\frac{p}{1-p}) + n \log(1-p)\} \quad (2)$$

- $\theta = p$
- $h(x) = \binom{n}{x}$
- $T(x) = x$
- $\eta(p) = \log(\frac{p}{1-p})$
- $A(p) = -n \log(1-p)$

Logistic regression and binary outcomes

Formulate model: Three elements

1. Error distribution for response variable
2. Linear predictor
3. Link function

The three elements of the logistic regression model are:

1. The Bernoulli probability distribution modelling the data: $P(y_i = 1 | x_i) = p_i$
2. The linear predictor: $\alpha + \sum_{j=1}^p \beta_j x_{ij}$
3. The link function g associating the mean of y , $P(y_i = 1 | x_i)$ to the linear predictor: here the link is the logistic link as we set $g(P(y_i = 1 | x_i)) = \text{logit}(p_i) = \beta_0 + \sum_{j=1}^p \beta_j x_{ij}$

glm() in R

- GLMs can be called in R just as linear models.

```
library(lars)
library(dplyr)

data(diabetes)
x <- as.data.frame.matrix(diabetes$x)
y <- ifelse(diabetes$y > mean(diabetes$y), 1, 0)
```

```
glm(y ~ age + sex + bmi + map + ltg, data = x, family = binomial) %>% summary()
```

Call:

```
glm(formula = y ~ age + sex + bmi + map + ltg, family = binomial,  
     data = x)
```

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-0.3164	0.1190	-2.659	0.00783	**
age	-1.0206	2.7309	-0.374	0.70860	
sex	-5.4254	2.6315	-2.062	0.03923	*
bmi	14.5079	3.0223	4.800	1.58e-06	***
map	11.8803	2.9652	4.007	6.16e-05	***
ltg	18.6940	3.1954	5.850	4.91e-09	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 606.61 on 441 degrees of freedom
Residual deviance: 434.17 on 436 degrees of freedom
AIC: 446.17

Number of Fisher Scoring iterations: 4

glm() in R

- Different types of exponential families can be called using the family option:
 - `binomial(link = 'logit')`
 - `gaussian(link = 'identity')`
 - `Gamma(link = 'inverse')`
 - `poisson(link='log')`
- There are similar return values as for the `lm` function:
 - coefficients
 - residuals

- fitted.values
- linear.predictors: the linear fit on the link scale

Making predictions

1. Train the prediction rule
2. Derive predictions on the linear predictor scale for the new data

```
library(lars)
library(dplyr)
library(ggplot2)
library(patchwork)
```

Warning: package 'patchwork' was built under R version 4.3.2

```
set.seed(11)

data(diabetes)
x <- as.data.frame.matrix(diabetes$x)
y <- ifelse(diabetes$y > mean(diabetes$y), 1, 0)

glm_predict <- glm(y ~ glu, data = x, family = binomial)
xnew <- data.frame(glu = rnorm(n = 1000, mean = 0, sd = 0.5))
xnew %>% head()
```

```
      glu
1 -0.29551555
2  0.01329718
3 -0.75827655
4 -0.68132667
5  0.58924458
6 -0.46707566
```

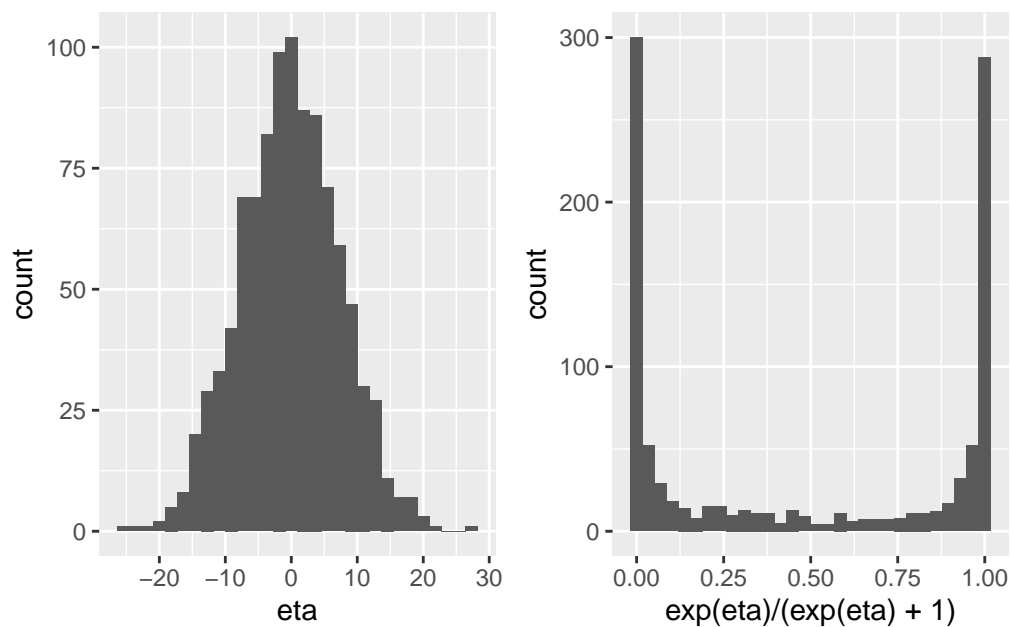
```
eta <- predict.glm(glm_predict, xnew)
```


Plot the predictions

```
ggplot() +  
  geom_histogram(aes(x = eta)) |  
ggplot() +  
  geom_histogram(aes(x = exp(eta) / (exp(eta) + 1)))
```

`stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

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```
rbinom(  
  n = length(eta),  
  size = rep(1, length(eta)),  
  prob = exp(eta) / (exp(eta) + 1)  
) %>% head()
```

```
[1] 0 0 0 0 1 0
```

Take away: Generalised linear model

The model formulation in GLMs consists of three elements:

1. Error distribution for response variable
2. Linear predictor
3. Link function

Most common data types can be modelled using GLMs

- Continuous \rightarrow Gaussian distribution
- Dichotomous or binary \rightarrow Bernoulli distribution
- Counts \rightarrow Poisson or Binomial (with known number of trial) distribution